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- A:n isolated nucleic acid molecule nucleic acid molecule comprising a polynucleotide having a nucleotide sequence at least 95% identical to a sequence selected from the group consisting of:
- (a) a nucleotide sequence encoding a full-length NAF-1 polypeptide having the complete amino acid sequence in SEQ ID NO:2, or the complete amino acid sequence encoded by the cDNA clone contained in the ATCC Deposit No. 97343;
- (b) a nucleotide sequence encoding a full-length NAF-1 polypeptide having the complete amino acid sequence in SEQ ID NO:2 excepting the N-terminal methionine (i.e., positions 1 to 331 of SEQ ID NO:2) or the complete amino acid sequence excepting the N-terminal methionine encoded by the cDNA clone contained in the ATCC Deposit No. 97343;
- (c) a nucleotide sequence encoding a predicted mature form of the NAF-1 polypeptide having the amino acid sequence at positions 24-331 or 27-331 in SEQ ID NO:2 or as encoded by the DNA cone contained in the ATCC Deposit No. 97343;
- (d) a nucleotide sequence discoding a polypeptide comprising the predicted TSR domain of the NAF-1 polypeptide in ving the amino acid sequence at positions 284-330 in SEQ ID NO:2 or as encoded by the cDNA clone contained in the ATCC Deposit No. 97343; and
- (e) a nucleotide sequence complementary to any of the nucleotide sequences in (a), (b), (c) or (d) above.
- The nucleic acid molecule of claim 1 wherein said polynucleotide has
 the complete nucleotide sequence in Figure 1 (SEQID NO:1).
 - 3. The nucleic acid molecule of claim 1 wherein said polynucleotide has the nucleotide sequence in Figure 1 (SEQ ID NO:1) encoding the NAF-1 polypeptide having the amino acid sequence in positions 2 to 331 of SEQID NO:2.

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The nucleic acid molecule of claim 1 wherein said polynucleotide has the nucleotide sequence in Figure 1 (SEQ ID NO:1) encoding the mature NAF-1 polypeptide having the amino acid sequence from about 27 to about 331 in SEQ ID NO:2.

5. At: solated nucleic acid molecule comprising a polynucleotide having a nucleotide sequence at least 95% identical to a sequence selected from the group consisting of:

(a) a nucleotide sequence encoding a polypeptide comprising the amino acid sequence of residues n-3 10f SEQ ID NO:2, where n is an integer in the range of 1-283;

(b) a nucleotide sequence encoding a polypeptide comprising the amino acid sequence of residues 1-m of SEQ ID NO:2, where m is either 330 or 331;

(c) a nucleotide sequence encoding a polypeptide having the amino acid sequence consisting of residues n-m of SEQ ID NO:2, where n and m are integers as defined respectively in (a) and (b) above; and

(d) a nucleotide sequence encoding a polypeptide consisting of a portion of the complete NAF-1 amino acid sequence encoded by the cDNA clone contained in ATCC Deport No. 97343 wherein said portion excludes from 1 to about 283 amino acids from the amino terminus of said complete amino acid sequence encoded by the cDNA clone contained in ATCC Deposit No. 97343;

(e) a nucleotide sequence encoding a polypeptide consisting of a portion of the complete NAF-1 amino acid sequence encoded by the cDNA clone contained in ATCC Deposit No. 97343 wherein said portion excludes 1 amino acid from the carboxy terminus of said complete amino acid sequence encoded by the cDNA clone contained in ATCC Deposit No. 97343; and

(f) a nucleotide sequence encoding a polypeptide consisting of a portion of the complete NAF-1 amino acid sequence encoded by the cDNA clone contained in ATCC Deposit No. 97343 wherein said portion include a combination of any of the amino terminal and carboxy terminal deletions in (d) and (e), above.

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- 6. The nucleic acid molecule of claim 1 wherein said polynucleotide has the complete nucleotide sequence of the cDNA clone contained in ATCC Deposit No. 97343.
- 7. The nucleic acid molecule of claim 1 wherein said polynucleotide has the nucleotide sequence encoding the NAF-1 polypeptide having the complete amino acid sequence excepting the N-terminal methionine encoded by the cDNA clone contained in ATCC/Deposit No. 97343.
 - 8. The nucleic acid molecule of claim 1 wherein said polynucleotide has the nucleotide sequence encoding the mature polypeptide having the amino acid sequence encoded by the cDNA clone contained in ATCC Deposit No. 97343.
 - 9. An isolated nucleic acid molecule comprising a polynucleotide which hybridizes under stringent hybridization conditions to a polynucleotide having a nucleotide sequence identical to a nucleotide sequence in (a), (b), (c), (d) or (e) of claim 1 wherein said polynucleotide which hybridizes does not hybridize under stringent hybridization conditions to a polynucleotide having a nucleotide sequence consisting of only A residues or of only T residues.
 - 10. An isolated nucleic acid molecule comprising a polynucleotide which encodes the amino acid sequence of an epitope-bearing portion of a NAF-1 polypeptide having an amino acid sequence in (a), (b), (c) or (d) of claim 1.
- 25 11. The isolated nucleic acid molecule of claim 10, which encodes an epitope-bearing portion of a NAF-1 polypeptide wherein the amino acid sequence of said portion is selected from the group of sequences in SEQ ID NO:2 consisting of: a polypeptide comprising amino acid residues from about Pro-75 to about Gly-100; a polypeptide comprising amino acid residues from about Th-168 to about Leu-180; a polypeptide comprising amino acid residues from about Asp 204 to about Ile-226; a polypeptide comprising amino acid residues from about Ile-288 to about Pro-281; and a polypeptide comprising amino acid residues from about Olu-291 to about Ser-327.

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- 2. A method for making a recombinant vector comprising inserting an isolated nucleic acid molecule of claim 1 into a vector.
 - 13. A recombinant vector produced by the method of claim 12.
- 14. A method of making a recombinant host cell comprising introducing the recombinant vector of claim 13 into a host cell.
 - 15. A recombinant host cell produced by the method of claim 14.
- 16. A recombinant method for producing a NAF-1 polypeptide, comprising culturing the recombinant host cell of claim 15 under conditions such that said polypeptide is expressed and recovering said polypeptide.
- 17. An isolated NAF- polypeptide comprising an amino acid sequence at least 95% identical to a sequence selected from the group consisting of:
- (a) the amino acid sequence of the full length NAF-1 polypeptide having the complete amino acid sequence shown in SEQ ID NO:2 or the complete amino acid sequence excepting the N-terminal methionine encoded by the cDNA clone contained in the ATCC Deposit No. 97343;
 - (b) the amino acid sequence of the full-length NAF-1 polypeptide having the complete amino acid sequence shown in SEQ ID NO:2 excepting the N-terminal methionine (i.e., positions 1-331 of SEQ ID NO:2) or the complete amino acid sequence excepting the N-terminal methionine encoded by the cDNA clone contained in the ATCC Deposit No. 97343;
 - (c) the amino acid sequence of the mature NAF-1 polypeptide having the amino acid sequence of residues 24-331 or 27-331 in SEQ ID NO:2, or the mature NAF-1 amino acid sequence as encoded by the cDNA clone contained in ATCC Deposit No. 97343; and

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(d) the amino acid sequence of the TSR domain of NAF-1 having the amino acid sequence of residues 284 to 330 of SEQ ID NO:2, or the amino acid sequence of the TSR domain of NAF-1 encoded by the cDNA clone contained in ATCC Deposit No. 97343.

Ar isolated polypeptide comprising an epitope-bearing portion of 18. the NAF-1 protein, wherein said portion is selected from the group consisting of: a polypeptide comprising amino acid residues from about Pro-75 to about Gly-100; a polypeptide compaising amino acid residues from about Thr-168 to about Leu-180; a polypeptide comprising amino acid residues from about Asp-204 to about Ile-226; a polypeptide comprising amino acid residues from about Ile-258 to about Pro-281; and a polypeptide comprising amino acid residues from about Glu-291 to about Ser-327.

An isolated antibody that binds specifically to a NAF-1 polypeptide 19. of claim 17.

An isolated nucleic adid molecule comprising a polynucleotide having 20. a sequence at leas. 95% identical to a sequence selected from the group consisting of:

the nucleotide sequence of clone HLHCE24R (shown as SEQ ID (a)

NO:15);

the nucleotide sequence of clone HLHDR83R (shown as SEQ ID (b) NO:16):

the nucleotide sequence of clone APTSB36R (shown as SEQ ID (c)

NO:17); 25

the nucleotide sequence of a portion of the sequence shown in Figure 1 (SEQ ID NO:1) wherein said portion comprises at least 50 contiguous nucleotides from nucleotide 1 to 650; and

a nucleotide sequence complementary to any of the nucleotide sequences in (a), (b), (c) and (d).